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## Amendments to the Claims

This listing of claims will replace all prior versions, and listings, of claims in the application:

## Listing of Claims

## 1-157. (canceled)

- 158. (Currently Amended) A method for identifying a protein target as being able to bind a ligand, comprising:
  - (a) providing a molecule comprising a methotrexate moiety which binds to dihydrofolate reductase, covalently bonded linked to the ligand, which methotrexate moiety binds to a dihydrofolate reductase;
  - introducing the molecule into a cell which i) (b) expresses a first fusion protein comprising a the dihydrofolate reductase capable of binding the methotrexate moiety, ii) expresses a second fusion protein comprising the protein target, wherein one of either the first and or second fusion protein proteins also comprises a transcription activator domain and the other fusion protein comprises a DNAbinding domain, and iii) has a reporter gene, reporter expression φf the wherein conditioned on the proximity of the first fusion protein to the second fusion protein and wherein the DNA-binding domain binds upstream of the reporter gene;

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- (c) permitting the molecule to bind to the first fusion protein and to the second fusion protein so as to activate the expression of the reporter gene; and
- (d) selecting the cell if it expresses the reporter gene,

so as to thereby identify the protein target as being able to bind the ligand.

- 159. (Currently Amended) The method of claim 158, wherein the protein target is encoded by a DNA from the group consisting of genomic DNA[[,]] or a cDNA and synthetic DNA.
- 160. (Previously Presented) The method of claim 158, wherein the ligand has a known biological function.
- 161. (Previously Presented) The method of claim 158, wherein the first fusion protein is (dihdrofolate reductase) -(DNA-binding domain).
- 162. (Previously Presented) The method of claim 158, wherein the first fusion protein is (dihdrofolate reductase) (LexA).
- 163. (Previously Presented) The method of claim 158, wherein the first fusion protein is (dihdrofolate reductase) (transcription activation domain).
- 164 (Previously Presented) The method of claim 158, wherein the first fusion protein is (dihdrofolate reductase)-(B42).

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- 165. (Previously Presented) The method of claim 158, wherein the second fusion comprises a DNA-binding domain.
- 166. (Previously Presented) The method of claim 158, wherein the second fusion protein comprises LexA.
- 167. (Previously Presented) The method of claim 158, wherein the second fusion protein comprises a transcription activation domain.
- 168. (Previously Presented) The method of claim 158, wherein the second fusion protein comprises B42.
- 169. (Currently Amended) The method of claim 158, wherein the cell is <del>S. cerevisiae or E. coli</del> <u>S. cerevisiae or E. coli</u>.
- 170. (Previously Presented) The method of claim 158, wherein the reporter gene is lacZ, Gal4 or Ura-3.
- 171. (Previously Presented) The method of claim 158, wherein the cell is a bacterial cell, the molecule comprises a methotrexate moiety bound to the ligand, the first fusion protein comprises a dihydrofolate reductase and a LexA, the second fusion protein comprises the protein target and B42, and the reporter gene is LacZ.
- 172. (Previously Presented) The method of claim 158, wherein the cell is a yeast cell, the molecule comprises a methotrexate moiety bound to the ligand, the first fusion protein comprises a dihydrofolate reductase and a LexA,

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the second fusion protein comprises the protein target and B42, and the reporter gene is Gal4.

173-176. (Cancelled)

- (Withdrawn Currently Amended) In a method 177. determining if a protein interacts with a ligand in vivo, wherein the method comprises activating a reporter gene by contacting a cell expressing two fusion proteins, the first comprising a dihydrofolate reductase ligand binding domain fused to a DNA-binding domain, which DNA-binding domain binds upstream of the reporter transcription comprising a second and the gene, activation domain fused to the protein, with a covalently linked hybrid-ligand so as to activate the reporter gene, the improvement comprising a covalently linked hybrid-ligand having a methotrexate moiety, which methotrexate moiety binds to the dihydrofolate reductase.
  - 178. (Withdrawn) The method of claim 177, wherein the DNAbinding domain is a LexA DNA-binding domain.
  - 179 (Withdrawn) The method of claim 177, wherein the transcription activation domain is B42.
  - 180 (Withdrawn) The method of claim 177, wherein the reporter gene is LacZ or Gal4.